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400 320 240 160 8 **GCTCCCTAATTAAACAACATTCAACTACTTTTTAGATCTATTTTTTACGTCTAGTTTTTACACAACTTAAAACTGTAATAA** TTTTGGTCTTTATTATTTTCTTTCTCTGACACTAAAATTACCTTTAGCACTCCTTTTCTTTTAAAATTAAAAA **ITCGCAGGTATTAAAGCCGACTTAAAACAAATGAGTGAAGAA** TCTAGATGTGTTCCGTTTAACTTTTTATCTATTTTAAAAGCGTCCATAATTTCGGCTGAATTTTGTTTACTCACTTCTT CITITCITITITATITATGIATAAAACICAATCATITICICITITITITITATITICIGGAGCCGAATIGICAGCT <u>AAAACCAGAAATAATAAAAAGAAGACTGTGATTTTTAATGGAAATCGTGAGGAAAAGAAAATTTTAATTTTCATTT</u> CGAGGGATTAATTTGTTGTAAGTTGATGAAAATCTAGATAAAAATGCAGATCAAAAATGTGTTGAATTTGACATTATT AGATCTACACAAGGCAAATTGAAAAAATAGATAAAAT

480 TGCGAGAGCAAAAAAACCCTTGCCGTTTTTTCAAATGACTTTGGAAAAATTCATTGTGAGCGGTAGCGAAACTTTGAA ACGCTCTCGTTTTTTGGGAACGGCAAAAAAGTTTACTGAAACCTTTTTTAAGTAACACTCGCCATCGCTTTGAAACTT

FIG._ 1A-1



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640 560 TAAAAAAATGTAACCTTTAAACTTTTTTTTTCCGTTTTCTTTGAGTTTACCTTTTTTTATAATAATATTTTTTCCTCTAGC Asn Phe Leu His Trp Lys Phe Glu Lys Ile Arg Gln Lys Lys Leu Lys Trp Lys Lys Tyr Tyr Lys Lys Arg Arg Ser

Asp Met Asp Phe Lys Ser Arg Lys Leu Thr Leu Asn Glu Lys Lys Asp Leu Glu Lys Ile Tyr Ala Glu Ser Glu Leu Lys

Ala Lys Lys Leu Gly Thr Gln Pro Gly Val Val Leu Glu Met Thr Met Lys Glu Met Met Lys Asn 11e Asn Leu Asp

800 TTAATGAAGAACAGCAGGTCAATATAGGAAATTATTCAAAAATAAAGTTGAGCATAGTAAATCAGATGATCTAGTAACG AATTACTTCTTTGTCGTCCAGTTATATCCTTTAATAAGTTTTTTATTTCAACTCGTATCATTTAGTCTACTAGATCATTGC

Val Asn Glu Glu Thr Ala Gly Gln Tyr Arg Lys Leu Phe Lys Asn Lys Val Glu His Ser Lys Ser Asp Asp Leu Val Thr

FIG._ 1A-2



GGACTATTAGAGTGTGGAACTCGAAATAGTTTTGATAAAACAAGAAGTGCCTTTCGTTTTTGTATTTGTGAGAATTCA

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1120 1040 880 960 Gly Leu Leu Glu Cys Gly Thr Arg Asn Ser Phe Asp Lys Thr Arg Ser Ala Phe Arg Phe Cys Ile Cys Glu Arg Ile Gln Ser Ala Ser Lys Arg Lys Thr Met Lys Glu. Ala Asp Thr Met Asp Asp. Ile. Phe Lys Arg Leu Lys Asn Asn Lys Ser Thr **TCTGCAAGTAAAAGAAAACAATGAAAGAAGCGGACACAATGGATGATATTTTTAAGAGGCTAAAAATAATAATCTAC** TTAACTCAAAACAAAACTATTCCTAAAAACTCACTTTTATCTTAAGTTACCTTACTATAAAGAGTGTTGTTTTTCTG GCAACTGAGAAAGAAGCTGATAATGCAAGAGAGTAAAAGATTTCGATACAATGAAAGCAAAAACTAAAGAGGCTTTTG **AATTGAGTTTTGTTTTGATAAGGATTTTTTGAGTGAAAATAGAATTCAATGGAATGATATTTCTCACAAAAAAAGAC** CGTTGACTCTTTCTTCGACTATTACGTTCTTCTCATTTTCTAAAGCTATGTTACTTTCGTTTTGATTTCTCCGAAAAC Glu Leu Ser Phe Val Phe Asp Lys Asp Phe Leu Ser Glu Asn Arg Ile Gln Trp Asn Asp Ile Ser His Asn Lys Lys Asp CCTGATAATCTCACACCTTGAGCTTTATCAAAACTATTTTGTTCTTCACGGAAAGCAAAAACATAAACACTCTCTTAAGT Gin Leu Arg Lys Glu Ala Asp Asn Ala Arg Arg Val Lys Asp Phe Asp Thr Met Lys Ala Lys Thr Lys Glu Ala Phe

1200 ATATGATCGTTATGCTGGATTCCTTTCTATTTGTTCGATTACAGGTTGCAGACCAGCAGAGGTTTTAAAGGGTATAGAGA TATACTAGCAATACGACCTAAGGAAAGATAAACAAGCTAATGTCCAACGTCTGGTCGTCTTCAAAATTTCCCATATCTCT

Tyr Asp Arg Tyr Ala Gly Phe Leu Ser IIe Cys Ser IIe Thr Gly Cys Arg Pro Ala Glu Val Leu Lys Gly IIe Glu

FIG._1B-1



Tyr Asp Arg Tyr Ala Gly Phe Leu Ser 11e Cys Ser 11e Thr Gly Cys Arg Pro Ala Glu Val Leu Lys Gly 11e Glu

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1280 1360 1440 Arg Thr Leu His Phe Asp Leu Ser Lys Tyr His Asp Asn Glu Gln Met Asn Tyr 11e Leu Ser Gln Leu Lys Asp Asn Lys **AGAACATTACATTTTGATTTATCAAAATATCATGATAATGAGCAAATGAATTATATTTTGTCGCAATTAAAAGATAATAA** TAAAAAGATGTTTGGTCTACCCTTCGAGATGTTGTCGAACTCCGTTATGGAGATGTAGGTTGTATCTTGCAAAAGTGACA TAGTAAGAAACAGATATGAGGATGGTATATCTTTTAAAATACTTGGTGCAAAGGTTGGAAATGACAGAGGGCAAAGCGAA <u> ATCATTCTTTGTCTATACTCCTACCATATAGAAATTTTATGAACCACGTTTCCAACCTTTACTGTCTCCGGTTTCGCTT</u> TCTTGTAATGTAAAACTAAATAGTTTTATAGTACTATTACTCGTTTACTTAATATAAAACAGCGTTAATTTTCTATTAT ATTTTTCTACAAACCAGATGGGAAGCTCTACAACAGCTTGAGGCAATACCTCTACATCCAACATAGAACGTTTTCACTGT Ile Val Arg Asn Arg Tyr Glu Asp Gly Ile Ser Phe Lys Ile Leu Gly Ala Lys Val Gly Asn Asp Arg Gly Gln Ser Glu

Tyr Thr Leu Arg His Arg Val Ala Ser Asp Leu Lys Ala Ser Gly Ala Asp Asp Phe Thr Ile Ala Ala ??? Leu Gly His

ATACACTTCGTCACAGGGTTGCGAGTGATCTCAAGGCATCCGGTGCAGATGACTTCACCATAGCGGCTNTTTTGGGTCAC

Phe Phe Tyr Lys Pro Asp Gly Lys Leu Tyr Asn Ser Leu Arg Gln Tyr Leu Tyr 11e Gln His Arg Thr Phe Ser Leu

TA T G T G A G C A G C C C C A A C C C T A G A G C C C A C G T C T C C A A G T G C C C C A A A A A A C C C A G T G

FIG._ 1B-2



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TCTCACTGAGTTTCGGTCCTCAATGATGCCGATACGAGCAGCAGCNTTCCACCATAGCGACATTGACCACAACTCACGA

1600

Arg Val Thr Gin Ser Gin Giu Leu Leu Arg Leu Cys Ser Phe Val ??? Arg Trp Tyr Arg Cys Asn Trp Cys

1660 CTGATGTTGTGAAAGCAAACAAGAGTCAGTTNGCTGTATCAAGGACTCCGAGCCAGATCT

GACTACAACACTTTCGTTTGTTCTCAGTCAANCGACATAGTTCCTGAGGCTCGGTCTAGA

FIG._1C



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160 240 320 400 80 **GCTGTTTTGGTCGAAACGTCCGACCGAAACTTCCTGATTTTCTTCACCCCTGAAGAAACTCAGAACATTAGTTCAACCAG AGAACTCGATTACGATTTGTAAGTAGAAATCTAACTCACATTTCGCAGAAAGTCAAACTTACCTCTTAGTTACAACTCAA** TGGGTTTTTGGGGTTGGATCAGAAAGTCGTTCAAGATTATTGACATAAAGACAGGAAGGTTTATAACAAGTACCAGATA **ACCCAAAAACCCCAACCTAGTCTTTTCAGCAAGTTCTAATAACTGTATTTCTGTCCTTCCAAATATTGTTCATGGTCTAT** AATCACCTATATAGACAATCGATGTCTCTTTTCGCTAAAATCTCCCAAACTGCTCCAAAAAGCTCGATAGGTCCCAAAA CGACAAAACCAGCTTTGCAGGCTGGCTTTGAAGGACTAAAAGAAGTGGGGACTTCTTTGAGTCTTGTAATCAAGTTGGTC TCTTGAGCTAATGCTAAACATTCATCTTTAGATTGAGTGTAAAGCGTCTTTCAGTTTGAATGGAGAATCAATGTTGAGTT TTAGTGGATATATCTGTTAGCTACAGAGAAAGCGATTTTAGAGGGTTTGACGAGGTTTTTCGAGCTATCCAGGGTTTT

<u> AAATTTCCTAGCCTTTTCAGATCCTTAAGCATACATATTTTGTTTAAACCGATTGTGTCCGGTGTTTGGTGTGGGAGCCAT</u>

TTTAAAGGATCGGAAAAGTCTAGGAATTCGTATGTATAAAACAAATTTGGCTAACACAGGCCACAAACCACACCTCGGTA



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720

560

640

800

TCTGTGGGTCAAGCCTCAAGTGCTGATCTGTGGTGTCTTCACCTGATAACTTTCACTTTTCGAGTGAAATTCAGGAGG

AGACACCCAGTTCGGAGTTCACGACTAGACACCACAGCAGATGGACTATTGAAAGTGAAAAAGCTCACTTTAAGTCCTCC

CGAAACGAGTGCCTAAGTGTTTTGGTTATCAACCAGGTAAGCTATGAGAAAGCCCAGCCATAAATGGGGTTAGGTTGAAG GTTCAGAAGTATACCACGCTGTGTTCCCCACATCATCCCAGAGTTTGACTTTCCAAACTATCGAGATTCGAACACGA GCTTTGCTCACGGATTCACAAAACCAATAGTTGGTCCATTCGATACTCTTTCGGGTCGGTATTTACCCCAATCCAACTTC CAAGTCTTCATATGGTGCGACACAAGGGGTGTAGTAGGGTGTCGTCAAACTGAAAGGTTTGATAGCTCTAAGCTTGTGCT

880 CGAAACTATGGGTCAAGCCCAGCTTTGCTGGGGTTCGGCACATCCAGCTTACAGCATTGGTGCTCTTGCGAAGCTGAAGC GCTTTGATACCCAGTTCGGGTCGAAACGACCCCAAGCCGTGTAGGTCGAATGTCGTAACCACGAGAACGCTTCGACTTCG

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1280

	ATTGATTTTTTTCTCCATTTATATACCTA	ATTCTCTATATA
00	AAGAGATATATCTATTAGCTTAAATATAACTAAAAAAGAGGTAAATATATGGAT	TITGACAGIGATTATTTAATATTAAGAGATATATCTATTAGCTTAAA
	TTCATACAATTTTTGCTCATTTTTGTTATTGAATATATAAATTAAGACTTAATAT	ATTATCTATATAGGCAATCTAATATTTCATACAATTTTTGCTCATTTTT
20	TAAAAACGAGTAAAACAATAACTTATATATTTAATTCTGAATTATA	AAGTATGI
		TITITATAAAACCTTAAAAATTTCCGCTATGAACGATGGCGTGAAAACGGTATAAATTTTGGACTGATAGAAATATTCA
040	ATATTTAAAACCTGACTATCTTTATAAGT	AAAAATATTTGGAATTTTTAAAGGCGATACTTGCTACCGCACTTTTGCCATATTTAAAACCTGACTATCTTTATAAGT
)	AAAAAATATGGTCTTCGTTTTGTTTTTTTTTTTTTTTTAAAAGCTCGCT	TGTTTTTAGATTAGGTCCCAAAACCCAAAAATATGGTCTTCGTTTGTTT
c	FACCAGAAGCAAAACAAAAAAAAAAAAAAAAAAAATTTTCGAGCGA	

1360 **GCATTACTTTGATATAGATAAAGCATTAAATGGTGATGATGTGGCGATATTATAAACTCAAATTTATCCATTGATGAAA** CGTAATGAAACTATATCTATTTCGTAATTTACCACTACTTACACCGCTATAATATTTGAGTTTAAATAGGTAACTACTTT



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1840



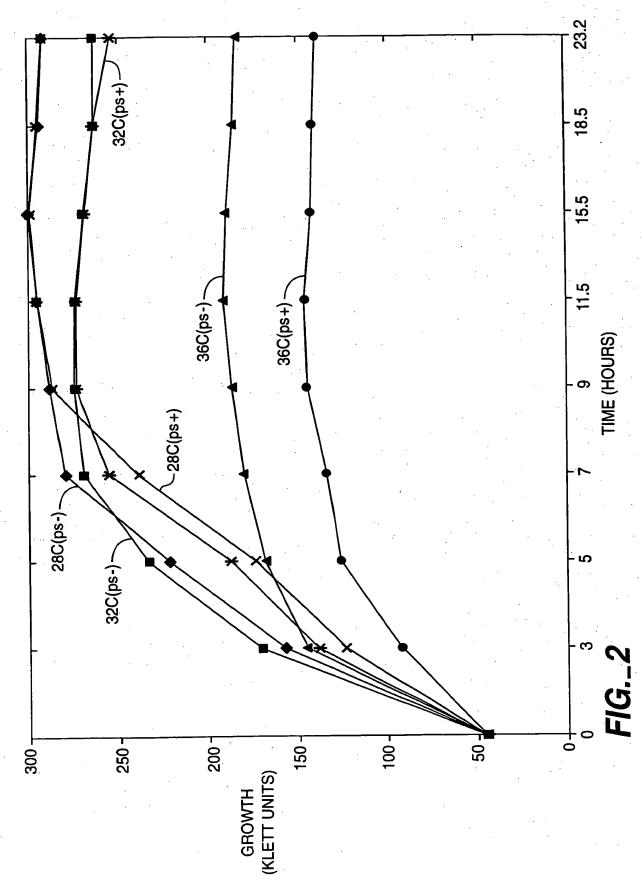
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